

Srivastava

#10/2400

1653

CRF Errors Corrected by the STIC Systems Branch

Serial Number: 09/424,487

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JUL 12 2000

SECRET 1600/2900

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically:  
\_\_\_\_\_
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by applicant was ☐ the prior application data; or ☐ other \_\_\_\_\_
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically:  
\_\_\_\_\_
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:  
\_\_\_\_\_
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:  
\_\_\_\_\_
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included:  
\_\_\_\_\_
- ☐ Deleted extra, invalid, headings used by an applicant, specifically:  
\_\_\_\_\_
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as \_\_\_\_\_
- ☒ Inserted mandatory headings, specifically: Seq 2
- ☐ Corrected an obvious error in the response, specifically:  
\_\_\_\_\_
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically:  
\_\_\_\_\_
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: \_\_\_\_\_
- ☒ Other: Seq 2 - corrected amino acid nos

RAW SEQUENCE LISTING                      DATE: 06/27/2000  
 PATENT APPLICATION: US/09/424,487        TIME: 21:14:19

Input Set : A:\Pto.amc  
 Output Set: N:\CRF3\06272000\I424487.raw

3 <110> APPLICANT: CHOO, Yen  
 4        KLUG, Aaron  
 5        ISALAN, Mark  
 7 <120> TITLE OF INVENTION: Nucleic Acid Binding Proteins  
 9 <130> FILE REFERENCE: 71278/264975  
 11 <140> CURRENT APPLICATION NUMBER: US 09/424,487  
 C--> 12 <141> CURRENT FILING DATE: 2000-02-29  
 14 <150> PRIOR APPLICATION NUMBER: GB 9710809.6  
 15 <151> PRIOR FILING DATE: 1997-05-23  
 17 <150> PRIOR APPLICATION NUMBER: PCT/GB98/01512  
 18 <151> PRIOR FILING DATE: 1998-05-26  
 20 <160> NUMBER OF SEQ ID NOS: 2  
 22 <170> SOFTWARE: PatentIn Ver. 2.1  
 24 <210> SEQ ID NO: 1  
 25 <211> LENGTH: 264  
 26 <212> TYPE: DNA  
 27 <213> ORGANISM: Artificial Sequence  
 29 <220> FEATURE:  
 30 <221> NAME/KEY: CDS  
 31 <222> LOCATION: (1)..(264)  
 33 <220> FEATURE:  
 34 <223> OTHER INFORMATION: Description of Artificial Sequence: encoding  
 35 <223> OTHER INFORMATION: nucleic acid binding proteins  
 37 <400> SEQUENCE: 1  
 38 gca gaa gag aag cct ttt cag tgt cga atc tgc atg cgt aac ttc agc    48  
 39 Ala Glu Glu Lys Pro Phe Gln Cys Arg Ile Cys Met Arg Asn Phe Ser  
 40    1                      5                      10                      15  
 42 gat cgt act act ctt acc cgc cac acg agg acc cac aca ggc gag aag    96  
 43 Asp Arg Thr Thr Leu Thr Arg His Thr Arg Thr His Thr Gly Glu Lys  
 44                      20                      25                      30  
 46 cct ttt cag tgt cga atc tgc atg cgt aac ttc agc agg agc gat aac    144  
 47 Pro Phe Gln Cys Arg Ile Cys Met Arg Asn Phe Ser Arg Ser Asp Asn  
 48                      35                      40                      45  
 50 ctt acg aga cac cta agg acc cac aca ggc gag aag cct ttt cag tgt    192  
 51 Leu Thr Arg His Leu Arg Thr His Thr Gly Glu Lys Pro Phe Gln Cys  
 52                      50                      55                      60  
 54 cga atc tgc atg cgt aac ttc agg caa gct gat cat ctt caa gag cac    240  
 55 Arg Ile Cys Met Arg Asn Phe Arg Gln Ala Asp His Leu Gln Glu His  
 56    65                      70                      75                      80  
 58 cta aag acc cac aca ggc gag aag                      264  
 59 Leu Lys Thr His Thr Gly Glu Lys  
 60                      85  
 63 <210> SEQ ID NO: 2  
 64 <211> LENGTH: 88  
 65 <212> TYPE: PRT  
 66 <213> ORGANISM: Artificial Sequence  
 W--> 67 <220> FEATURE:

RAW SEQUENCE LISTING                      DATE: 06/27/2000  
 PATENT APPLICATION: US/09/424,487        TIME: 21:14:19

Input Set : A:\Pto.amc  
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68 <223> OTHER INFORMATION: Description of Artificial Sequence: encoding  
 69 <223> OTHER INFORMATION: nucleic acid binding proteins  
 71 <400> SEQUENCE: 2  
 72 Ala Glu Glu Lys Pro Phe Gln Cys Arg Ile Cys Met Arg Asn Phe Ser  
 73 1                      5                      10                      15  
 75 Asp Arg Thr Thr Leu Thr Arg His Thr Arg Thr His Thr Gly Glu Lys  
 76                      20                      25                      30  
 78 Pro Phe Gln Cys Arg Ile Cys Met Arg Asn Phe Ser Arg Ser Asp Asn  
 79                      35                      40                      45  
 81 Leu Thr Arg His Leu Arg Thr His Thr Gly Glu Lys Pro Phe Gln Cys  
 82                      50                      55                      60  
 84 Arg Ile Cys Met Arg Asn Phe Arg Gln Ala Asp His Leu Gln Glu His  
 85 65                      70                      75                      80  
 87 Leu Lys Thr His Thr Gly Glu Lys  
 88                      85

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VERIFICATION SUMMARY                      DATE: 06/27/2000  
PATENT APPLICATION:    US/09/424,487        TIME: 21:14:20

Input Set : A:\Pto.amc  
Output Set: N:\CRF3\06272000\I424487.raw

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:67 M:283 W: Missing Blank Line separator, <220> field identifier